```
Sequence Listing could not be accepted due to errors.
See attached Validation Report.
If you need help call the Patent Electronic Business Center at (866)
217-9197 (toll free).
Reviewer: Durreshwar Anjum
Timestamp: [year=2007; month=12; day=13; hr=10; min=18; sec=48; ms=335;
Reviewer Comments:
<210> 21
<211> 21
<212> DNA
<213> Artificial Sequence
<220>
<221> misc DNA
<222> (20)..(21)
<223> n stands for deoxy thymidine
<400> 21
                                     21
ggaccaggaa auuccgauun n
```

If <213> response is Artificial, please give the source of genetic material in numeric identifier <223>, this type of error is shown in seq id 22.

Validated By CRFValidator v 1.0.3

Application No: 10542408 Version No: 1.0

Input Set:

Output Set:

Started: 2007-11-21 18:01:39.451 **Finished:** 2007-11-21 18:01:41.039

Elapsed: 0 hr(s) 0 min(s) 1 sec(s) 588 ms

Total Warnings: 16
Total Errors: 4

No. of SeqIDs Defined: 22

Actual SeqID Count: 22

Error code		Error Description
W	213	Artificial or Unknown found in <213> in SEQ ID (5)
W	213	Artificial or Unknown found in <213> in SEQ ID (6)
W	213	Artificial or Unknown found in <213> in SEQ ID (7)
W	213	Artificial or Unknown found in <213> in SEQ ID (10)
W	213	Artificial or Unknown found in <213> in SEQ ID (11)
W	213	Artificial or Unknown found in <213> in SEQ ID (12)
M	213	Artificial or Unknown found in <213> in SEQ ID (13)
W	213	Artificial or Unknown found in <213> in SEQ ID (14)
W	213	Artificial or Unknown found in <213> in SEQ ID (15)
W	213	Artificial or Unknown found in <213> in SEQ ID (16)
W	213	Artificial or Unknown found in <213> in SEQ ID (17)
W	213	Artificial or Unknown found in <213> in SEQ ID (18)
W	213	Artificial or Unknown found in <213> in SEQ ID (19)
W	213	Artificial or Unknown found in <213> in SEQ ID (20)
W	213	Artificial or Unknown found in <213> in SEQ ID (21)
E	257	Invalid sequence data feature in <221> in SEQ ID (21)
E	224	$<\!220\!>, <\!223\!>$ section required as $<\!213\!>$ has Artificial sequence or Unknown in SEQID (21)
W	213	Artificial or Unknown found in <213> in SEQ ID (22)
E	257	Invalid sequence data feature in <221> in SEQ ID (22)

Input Set:

Output Set:

Started: 2007-11-21 18:01:39.451 **Finished:** 2007-11-21 18:01:41.039

Elapsed: 0 hr(s) 0 min(s) 1 sec(s) 588 ms

Total Warnings: 16
Total Errors: 4
No. of SeqIDs Defined: 22

Actual SeqID Count: 22

Error code Error Description

E 224 <220>,<223> section required as <213> has Artificial sequence or

Unknown in SEQID (22)

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Gly Asp His Arg Leu Val Leu Ala Ala Val Glu Thr Thr Val Leu Val
                           40
Leu Ile Phe Ala Val Ser Leu Leu Gly Asn Val Cys Ala Leu Val Leu
                        55
Val Ala Arg Arg Arg Arg Gly Ala Thr Ala Cys Leu Val Leu Asn
                   70
                                      75
Leu Phe Cys Ala Asp Leu Leu Phe Ile Ser Ala Ile Pro Leu Val Leu
Ala Val Arg Trp Thr Glu Ala Trp Leu Leu Gly Pro Val Ala Cys His
                              105
           100
Leu Leu Phe Tyr Val Met Thr Leu Ser Gly Ser Val Thr Ile Leu Thr
                          120
                                              125
Leu Ala Ala Val Ser Leu Glu Arg Met Val Cys Ile Val His Leu Gln
                       135
Arg Gly Val Arg Gly Pro Gly Arg Arg Ala Arg Ala Val Leu Leu Ala
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                                     155
Leu Ile Trp Gly Tyr Ser Ala Val Ala Ala Leu Pro Leu Cys Val Phe
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Phe Arg Val Val Pro Gln Arg Leu Pro Gly Ala Asp Gln Glu Ile Ser

180 185 Ile Cys Thr Leu Ile Trp Pro Thr Ile Pro Gly Glu Ile Ser Trp Asp 200 Val Ser Phe Val Thr Leu Asn Phe Leu Val Pro Gly Leu Val Ile Val 215 220 Ile Ser Tyr Ser Lys Ile Leu Gln Ile Thr Lys Ala Ser Arg Lys Arg 235 230 Leu Thr Val Ser Leu Ala Tyr Ser Glu Ser His Gln Ile Arg Val Ser 245 250 Gln Gln Asp Phe Arg Leu Phe Arg Thr Leu Phe Leu Leu Met Val Ser 260 265 270 Phe Phe Ile Met Trp Ser Pro Ile Ile Ile Thr Ile Leu Leu Ile Leu 275 280 285 Ile Gln Asn Phe Lys Gln Asp Leu Val Ile Trp Pro Ser Leu Phe Phe 295 300 Trp Val Val Ala Phe Thr Phe Ala Asn Ser Ala Leu Asn Pro Ile Leu 310 315 305 Tyr Asn Met Thr Leu Cys Arg Asn Glu Trp Lys Lys Ile Phe Cys Cys 325 330 Phe Trp Phe Pro Glu Lys Gly Ala Ile Leu Thr Asp Thr Ser Val Lys 345 350 Arg Asn Asp Leu Ser Ile Ile Ser Gly 355 360

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20 25 30

Gly Asp His Arg Leu Val Leu Ser Val Val Glu Thr Thr Val Leu Gly
35 40 45

Leu Ile Phe Val Val Ser Leu Leu Gly Asn Val Cys Ala Leu Val Leu 50 55 60

Val Ala Arg Arg Arg Arg Gly Ala Thr Ala Ser Leu Val Leu Asn
65 70 75 80

Leu Phe Cys Ala Asp Leu Leu Phe Thr Ser Ala Ile Pro Leu Val Leu
85 90 95

Val Val Arg Trp Thr Glu Ala Trp Leu Leu Gly Pro Val Val Cys His
100 105 110

Leu Leu Phe Tyr Val Met Thr Met Ser Gly Ser Val Thr Ile Leu Thr
115 120 125

Leu Ala Ala Val Ser Leu Glu Arg Met Val Cys Ile Val Arg Leu Arg 130 135 140

Arg Gly Leu Ser Gly Pro Gly Arg Arg Thr Gln Ala Ala Leu Leu Ala 145 150 155 160

Phe Ile Trp Gly Tyr Ser Ala Leu Ala Ala Leu Pro Leu Cys Ile Leu 165 170 175

Phe Arg Val Val Pro Gln Arg Leu Pro Gly Gly Asp Gln Glu Ile Pro 180 185 190

Ile Cys Thr Leu Asp Trp Pro Asn Arg Ile Gly Glu Ile Ser Trp Asp 195 200 205

Val Phe Phe Val Thr Leu Asn Phe Leu Val Pro Gly Leu Val Ile Val
210 215 220

Ile Ser Tyr Ser Lys Ile Leu Gln Ile Thr Lys Ala Ser Arg Lys Arg

Leu Thr Leu Ser Leu Ala Tyr Ser Glu Ser His Gln Ile Arg Val Ser

245 250 255

Gln Gln Asp Tyr Arg Leu Phe Arg Thr Leu Phe Leu Leu Met Val Ser

Phe Phe Ile Met Trp Ser Pro Ile Ile Ile Thr Ile Leu Leu Ile Leu 275 280 285

Ile Gln Asn Phe Arg Gln Asp Leu Val Ile Trp Pro Ser Leu Phe Phe 290 295 300

Trp Val Val Ala Phe Thr Phe Ala Asn Ser Ala Leu Asn Pro Ile Leu 305 310 315 320

Tyr Asn Met Ser Leu Phe Arg Asn Glu Trp Arg Lys Ile Phe Cys Cys
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gtegtggaga ceaeegttet ggggeteate tttgtegtet eaetggtgg eaaeggtgt 180
getetagtge tggtggegeg eegteggege egtgggega eageeageet ggtgeteaae 240
etettetgeg eggatttget etteaeeage geeateeete tagtgetegt egtgegetgg 300

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actgaggeet ggetgttggg geeegtegte tgeeacetge tettetaegt gatgaeaatg
                                                                   360
ageggeageg teaegateet caeaetggee geggteagee tggagegeat ggtgtgeate
                                                                    420
gtgcgcctcc ggcgcggctt gagcggcccg gggcggcgga ctcaggcggc actgctggct
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ttcatatggg gttactcggc gctcgccgcg ctgcccctct gcatcttgtt ccgcgtggtc
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ccgcagcgcc ttcccggcgg ggaccaggaa attccgattt gcacattgga ttggcccaac
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cgcataggag aaatctcatg ggatgtgttt tttgtgactt tgaacttcct ggtgccggga
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                                                                    720
ctggtcattg tgatcagtta ctccaaaatt ttacagatca cgaaagcatc gcggaagagg
cttacgctga gcttggcata ctctgagagc caccagatcc gagtgtccca acaagactac
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cgactcttcc gcacgctctt cctgctcatg gtttccttct tcatcatgtg gagtcccatc
                                                                    840
atcatcacca tectectcat ettgatecaa aactteegge aggaeetggt catetggeea
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tecettttet tetgggtggt ggeetteaeg tttgeeaaet etgeeetaaa eeceataetg
tacaacatgt cgctgttcag gaacgaatgg aggaagattt tttgctgctt cttttttcca 1020
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ccgcagcgcc ttcccggcgg ggaccaggaa attccgattt gcacattgga ttggcccaac
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cgcataggag aaatctcatg ggatgtgttt tttgtgactt tgaacttcct ggtaccagga
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ctggtcattg tgatcagcta ctccaagatt ttacagatca cgaaagcctc gcggaagagg
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cttacgctga gcttggcata ctccgagagc caccagatcc gagtgtccca gcaggactac
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eggetettee gaaegetett eetgeteatg gttteettet teateatgtg gagteeeate
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atcatcacca tectectcat ettgatecag aactteegge aggaeetggt tatetggeeg
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tcccttttct tctgggtggt ggccttcacg tttgccaact ccgccctaaa ccccattctg
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tacaacatgt cgctgttcag gagcgagtgg aggaagattt tttgctgctt ctttttccca
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caactccgcc ctaaacccca ttctgt
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<223> n stands for deoxy thymidine
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ggaccaggaa auuccgauun n
                                          21
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